SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: Mundy, Gregory R.
Gallwitz, Wolfgang E.

(ii) TITLE OF INVENTION: SCREENING ASSAY FOR THE IDENTIFICATION OF AGENTS WHICH INHIBIT CANCER METASTASIS TO BONE

(iii) NUMBER OF SEQUENCES: 1

(iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: Arnold, White & Durkee
(B) STREET: P.O. Box 4433
(C) CITY: Houston
(D) STATE: Texas
(E) COUNTRY: USA
(F) ZIP: 77210

(v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER: US Unknown
(B) FILING DATE: Concurrently Herewith
(C) CLASSIFICATION: Unknown

(viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Highlander, Steven L.
(B) REGISTRATION NUMBER: 37,642
(C) REFERENCE/DOCKET NUMBER: OSTS:002PZ1

(ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: 512/418-3000
(B) TELEFAX: 512/474-7577

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 4348 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

AGGATCCCAT CAGCTTGATG CATATCTATA CACTCCTCCC TGAGGCAGTT CCTCCAGAGG 60

TTAGCAGCCC GCCCTGTTCC TGGAGAAGTC TTATCCTCAC CTAACTACAA AAAGCATTTA 120

ATAAAGAAAC ACACCCTTTC CTATTGTTAC TTGGGGTTTG AAGGCATTAC ATCTTTTTTT 180

CTTTTTTGCT ACCTTGAGGT CAGCTGGCAA CAGCCTCCTC TCAAGTCTCA GTCCAGGCGA 240

GCGATGAGAG CCACATTCTA ATGGAATCCA ATGAGAGCAT TACTTGAGCT GATTATGCAA 300

CGTCTCTTTG TAAACATTGA AAAAAGTTTG GAGAAGAGAT GGGATGAGTT TAGTTGTTTG 360

GTTCCAGGGG ATTTTAGAGA CATACGTTGC AGCTACAGAT TGGTAAATGT GAAATCTGGA 420

TGCTTATTGG TAATAAGAGA ATTCCCAAGC CCAGGTGCAC TGTTTAAAGT GCTATAGATT 480

CATATTTGGT TTATAATGTA TATCTGTTTG CTGTTTGGGT TAAGGAGGAA AGAAGAAAGA 540

AGGATGAAGA GGCTAAGCAT AAATGCTATT TACTTTTTTC TAAGCTATGA CAGGAGATAT 600

ACATTAACTG GGTATTCAAC TGAATTTAAG AGTAATGCAT TTAAAATTTT TTCAACCTCT 660

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CTTGGGAGAT GCCCTTGATG TTGAAGAGGC TTTTTGAGAG CGGGCTAAAA AGGGGGAGCG 4260

GAGTAGTGCG GGGAGATGGA GAGTCCTGAC TGACACCTCG GGTCCCATTC CCTTCTGTTG 4320

CAGGTCCCGA GCGCGAGCGG AGACGATG 4348